

## SEQUENCE LISTING

<110> Alexandra Charlesworth
 Falvia Spirito
 Guerrino Meneguzzi
 John Baird
 Keith Linder

<120> ISOLATION OF THE LAMININ Y2 GENE IN HORSES AND ITS USE IN DIAGNOSING JUNCTIONAL EPIDERMOLYSIS BULLOSA

<130> p84us4

<160> 32

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 3989

<212> DNA

<213> equine

<220>

<221> CDS

<222> (198) . . . (3767)

<223>

<400> 1

ttattcacag	gtgagtcaca	ccctgaaaca	caggetetet	tcctgtcagg	60
gtagaagagt	cgataaaacc	acctgatcaa	ggaaaaggaa	ggcacagcgg	120
gagaactccc	agcggcgagg	cgccgggcag	gegaeceetge	agcggcggac	180
cctggcc ato	cct gcg c	tc tgg ctg	agc tgc tac	ctc tgc	230
Met	: Pro Ala I	eu Trp Leu	Ser Cys Tyr	Leu Cys	
1		5		10	
	gtagaagagt gagaactccc cctggcc atg	gtagaagagt cgataaaacc gagaactccc agcggcgagg cctggcc atg cct gcg c	gtagaagagt cgataaaacc acctgatcaa gagaactccc agcggcgagg cgccgggcag cctggcc atg cct gcg ctc tgg ctg	gtagaagagt cgataaaacc acctgatcaa ggaaaaggaa gagaactccc agcggcgagg cgccgggcag cgacccctgc cctggcc atg cct gcg ctc tgg ctg agc tgc tac	tattcacag gtgagtcaca ccctgaaaca caggetetet tectgtcagg gtagaagagt cgataaaacc acctgatcaa ggaaaaggaa ggcacagegg gagaactece ageggegag egeegggeag egacecetge ageggeggac eetggee atg eet geg ete tgg etg age tge tae ete tge  Met Pro Ala Leu Trp Leu Ser Cys Tyr Leu Cys  1 5 10

ttc tcg ctc ctc ctg ccc gca gcc cgg gcc acc tcc ggg agg gaa gtc

Phe Ser Leu Leu Pro Ala Ala Arg Ala Thr Ser Gly Arg Glu Val

15 20 25

tgt gat tgc aac ggg aag tcc agg caa tgc atc ttt gac cag gaa ctt

Cys Asp Cys Asn Gly Lys Ser Arg Gln Cys Ile Phe Asp Gln Glu Leu

cac aaa cag aca gga aat gga ttc cgc tgc ctc aac tgc aat gac aac
His Lys Gln Thr Gly Asn Gly Phe Arg Cys Leu Asn Cys Asn Asp Asn
45
50
55

act gat ggc atc cac tgc gag agg tgc aag gca gga ttt tac cga cag

Thr Asp Gly Ile His Cys Glu Arg Cys Lys Ala Gly Phe Tyr Arg Gln

70

75

aga gaa agg gac cgc tgt tta ccc tgc aat tgt aac tct aaa ggt tct 470
Arg Glu Arg Asp Arg Cys Leu Pro Cys Asn Cys Asn Ser Lys Gly Ser
80 85 90

1				•														
 ٠			•															
							tct Ser									518	3	
gtg Val	aca Thr	gga Gly 110	gac Asp	agg Arg	tgt Cys	gac Asp	cga Arg 115	tgt Cys	ctg Leu	ccc Pro	ggc Gly	ttc Phe 120	cac His	aca Thr	ctc Leu	566	5	
act Thr	gat Asp 125	gct Ala	ggg Gly	tgc Cys	gcc Ala	caa Gln 130	gac Asp	caa Gln	agg Arg	ctg Leu	cta Leu 135	gac Asp	tcc Ser	aag Lys	tgt Cys	614	Į.	
gac Asp 140	tgt Cys	gac Asp	cca Pro	gct Ala	ggc Gly 145	atc Ile	tca Ser	gly ggg	ccc Pro	tgt Cys 150	gac Asp	tca Ser	ggc Gly	cgc Arg	tgt Cys 155	662	2	
							gga Gly									710	)	
ggt Gly	tac Tyr	tat Tyr	cac His 175	ctg Leu	gat Asp	gly ggg	gga Gly	aac Asn 180	cct Pro	cag Gln	ggc Gly	tgt Cys	acc Thr 185	cag Gln	tgt Cys	758	3	
ttt Phe	tgc Cys	tat Tyr 190	gly ggg	cat His	tcc Ser	gcc Ala	agc Ser 195	tgc Cys	cac His	agc Ser	tct Ser	999 Gly 200	gac Asp	tac Tyr	agt Ser	806		
							ttc Phe									854	ł	
gct Ala 220	gtc Val	caa Gln	aga Arg	aac Asn	999 Gly 225	tct Ser	cct Pro	gca Ala	aag Lys	ctc Leu 230	cag Gln	tgg Trp	tca Ser	cag Gln	cgc Arg 235	902	2	
							gca Ala									950	)	
							ggg Gly									998	3	
							gtg Val 275									1046	5	
gcc Ala	cat His 285	gac Asp	gtg Val	atc Ile	ctg Leu	gaa Glu 290	ggt Gly	gct Ala	ggt Gly	cta Leu	cgg Arg 295	atc Ile	aca Thr	gct Ala	ccc Pro	1094	1	
ttg Leu 300	Met	cca Pro	ctt Leu	agc Ser	aag Lys 305	aca Thr	ctg Leu	cct Pro	tgt Cys	999 Gly 310	atc Ile	acc Thr	aag Lys	act Thr	tac Tyr 315	1142	2	

ř n	•	,			•												
					aat Asn 320												1190
	agt Ser	tac Tyr	ttt Phe	gag Glu 335	tat Tyr	cgg Arg	agg Arg	tta Leu	ctg Leu 340	cgg Arg	aac Asn	ctc Leu	aca Thr	gcc Ala 345	ctg Leu	cgg Arg	1238
	atc Ile	cga Arg	gct Ala 350	acc Thr	tac Tyr	gga Gly	gaa Glu	tac Tyr 355	agt Ser	act Thr	ggg Gly	tac Tyr	att Ile 360	gac Asp	aac Asn	gtg Val	1286
					gcc Ala												1334
					tgc Cys												1382
	tgt Cys	gct Ala	tcc Ser	ggc Gly	tac Tyr 400	aaa Lys	aga Arg	gat Asp	tca Ser	gcc Ala 405	aga Arg	ctg Leu	gga Gly	cct Pro	ttt Phe 410	ggc Gly	1430
					tgt Cys												1478
	aca Thr	gga Gly	gac Asp 430	tgt Cys	tac Tyr	tca Ser	ggg Gly	gat Asp 435	gag Glu	aac Asn	cct Pro	gac Asp	atc Ile 440	cct Pro	gag Glu	tgt Cys	1526
	gct Ala	gac Asp 445	tgc Cys	ccc Pro	att Ile	ggt Gly	ttc Phe 450	tac Tyr	aac Asn	gat Asp	cca Pro	caa Gln 455	gac Asp	ccc Pro	cgc Arg	agc Ser	1574
					ccc Pro												1622
	gag Glu	aca Thr	gag Glu	gag Glu	gtg Val 480	gtg Val	tgc Cys	aat Asn	aac Asn	tgc Cys 485	ccc Pro	cag Gln	ggt Gly	gtc Val	act Thr 490	ggt Gly	1670
	gcc Ala	cgc Arg	tgt Cys	gag Glu 495	ctc Leu	tgt Cys	gct Ala	gat Asp	ggc Gly 500	tat Tyr	ttt Phe	gly 999	gac Asp	ccc Pro 505	ttc Phe	gly aaa	1718
					gtg Val												1766
					gcc Ala												1814
	ctg	aag	tgc	atc	cac	aac	aca	gct	<b>3</b> 99	gtc	cac	tgt	gac	cag	tgc	aaa	1862

•	7				•												
•	Leu 540	Lys	Cys	Ile	His	Asn 545	Thr	Ala	Gly	Val	His 550	Cys	Asp	Gln	Cys	Lys 555	
	gca Ala	ggc Gly	tac Tyr	tat Tyr	ggg Gly 560	gac Asp	ccg Pro	ttg Leu	gct Ala	ccc Pro 565	aat Asn	cca Pro	gca Ala	gac Asp	aag Lys 570	tgt Cys	1910
														gag Glu 585			1958
														ctc Leu			2006
	gag Glu	cat His 605	gcg Ala	gca Ala	ctg Leu	acc Thr	agc Ser 610	tgt Cys	cca Pro	gct Ala	tgc Cys	tat Tyr 615	aat Asn	caa Gln	gtg Val	aag Lys	2054
	gtt Val 620	cag Gln	atg Met	gat Asp	cag Gln	ttt Phe 625	atg Met	cag Gln	cag Gln	ctc Leu	cag Gln 630	atc Ile	ctg Leu	gag Glu	gcc Ala	ctg Leu 635	2102
	att Ile	tcg Ser	aag Lys	gct Ala	cag Gln 640	ggt Gly	gga Gly	gca Ala	gta Val	ccc Pro 645	aac Asn	gca Ala	gag Glu	ctg Leu	gaa Glu 650	ggc Gly	2150
	agg Arg	atg Met	cag Gln	cag Gln 655	gct Ala	gag Glu	cag Gln	gcc Ala	ctt Leu 660	cgg Arg	gac Asp	att Ile	ctg Leu	aga Arg 665	gaa Glu	gcc Ala	2198
	cag Gln	Ile	Ser	Gln	Asp	Ala	gtt Val	Arg	Ser	Phe	Asn	Leu	Arg	gtg Val	gcc Ala	aag Lys	2246
	gca Ala	agg Arg 685	act Thr	caa Gln	gag Glu	aat Asn	agc Ser 690	tac Tyr	cgg Arg	gac Asp	cgc Arg	ctg Leu 695	gat Asp	gac Asp	ctc Leu	aag Lys	2294
	atg Met 700	act Thr	gtg Val	gaa Glu	aga Arg	gtt Val 705	cgg Arg	gcc Ala	ctg Leu	ggc Gly	agt Ser 710	cag Gln	tat Tyr	cag Gln	aac Asn	caa Gln 715	2342
	gtt Val	cag Gln	gat Asp	act Thr	cgc Arg 720	agg Arg	ctc Leu	atc Ile	act Thr	cag Gln 725	atg Met	cgc Arg	ctg Leu	agc Ser	ctg Leu 730	gag Glu	2390
														tca Ser 745			2438
														gcc Ala			2486
														caa Gln			2534
												•					

765 770 775

													gtg Val			2582
													gcc Ala			2630
caa Gln	agg Arg	ctt Leu	gtg Val 815	gga Gly	aaa Lys	ttg Leu	cag Gln	aaa Lys 820	act Thr	aaa Lys	tct Ser	ctg Leu	gcc Ala 825	cag Gln	gag Glu	2678
ttg Leu	tcg Ser	agg Arg 830	gag Glu	gcc Ala	acg Thr	caa Gln	acc Thr 835	gac Asp	atg Met	gaa Glu	gca Ala	gat Asp 840	agg Arg	tct Ser	tat Tyr	2726
													cag Gln			2774
													caa Gln			2822
gat Asp	tct Ser	ctc Leu	tca Ser	aac Asn 880	cgt Arg	gtg Val	act Thr	aag Lys	cat His 885	atg Met	gat Asp	gag Glu	ttc Phe	aag Lys 890	cac His	2870
gtg Val	caa Gln	agc Ser	aat Asn 895	ctg Leu	gga Gly	aac Asn	tgg Trp	gaa Glu 900	gaa Glu	gaa Glu	acc Thr	cgg Arg	cag Gln 905	ctc Leu	tta Leu	2918
cag Gln	aat Asn	gga Gly 910	aag Lys	aat Asn	Gly 999	aga Arg	cag Gln 915	aca Thr	tca Ser	gat Asp	cag Gln	ctg Leu 920	ctt Leu	tcc Ser	cgt Arg	2966
													atg Met			3014
													aga Arg			3062
gac Asp	ctg Leu	cag Gln	gtt Val	gga Gly 960	gat Asp	aaa Lys	aga Arg	gca Ala	gaa Glu 965	gct Ala	gaa Glu	gag Glu	gcc Ala	atg Met 970	aag Lys	3110
													gac Asp 985			3158
			Glu					Ser					gcc Ala O			3206

gca aag aat gca Ala Lys Asn Ala 1005		Ala Leu G		Gly Lys		3254
cag gag ata gga Gln Glu Ile Gly 1020						3302
gcc ttg gcc atg Ala Leu Ala Met	gag aag gga Glu Lys Gly 1040	Leu Ala T	ct ctg aaa Thr Leu Lys .045	Ser Glu	atg aga Met Arg 1050	3350
gaa gtg gaa gga Glu Val Glu Gly 1055	Glu Leu Ser	agg aag g Arg Lys G 1060	gag cag gag Slu Gln Glu	ttt gac Phe Asp 1065	Met Asp	3398
atg gac gca gtg Met Asp Ala Val 1070						3446
aga gcc aag aat Arg Ala Lys Asn 1085	gct gga gtt Ala Gly Val 109	Thr Ile G	aa gac aca In Asp Thr 1095	Leu Asn	aca ttg Thr Leu	3494
gat ggc atc cta Asp Gly Ile Leu 1100						3542
agg ctg atc tta Arg Leu Ile Leu	ctg gag cag Leu Glu Gln 1120	Lys Leu P	tc cga gcc Phe Arg Ala .125	aag act Lys Thr	cag atc Gln Ile 1130	3590
aac agc cag cta Asn Ser Gln Leu 1135	Arg Pro Leu				Ala His	3638
cgg cag aag ggc Arg Gln Lys Gly 1150						3686
ctg gct gat gtg Leu Ala Asp Val 1165	aag aac ctg Lys Asn Leu 117	Glu Asn I	itc agg gac le Arg Asp 1175	Asn Leu	ccc ccg Pro Pro	3734
ggc tgc tac aat Gly Cys Tyr Asn 1180				agctgcc t	tagagattt	3787
ctcaaccaag gttch ttgggattca gacch tgaatatgtt gaato ggccagataa atgto	caggg ctcag gcgtgt gctca	gagcc cgca	itgcggg tggg	ggtggga t	gggaatatt	3847 3907 3967 3989

<210> 2 <211> 1190

<212> PRT

375

Pro Val Gly Tyr Lys Gly Gln Phe Cys Gln Asp Cys Ala Ser Gly Tyr

Lys Arg Asp Ser Ala Arg Leu Gly Pro Phe Gly Thr Cys Ile Pro Cys

380

395

	0				•											
				•	405					410					415	
•	Asn	Cys	Gln	-	405 Gly	Gly	Ala	Cys		410 Pro	Asp	Thr	Gly	Asp	415 Cys	Tyr
	Ser	Gly	_	420 Glu	Asn	Pro	Asp		425 Pro	Glu	Cys	Ala		430 Cys	Pro	Ile
	Gly		435 Tyr	Asn	Asp	Pro		440 Asp	Pro	Arg	Ser		445 Lys	Pro	Cys	Pro
		450 Arg	Asn	Gly	Phe		455 Cys	Ser	Val	Met		460 Glu	Thr	Glu	Glu	
	465 Val	Cys	Asn	Asn		470 Pro	Gln	Gly	Val		475 Gly	Ala	Arg	Cys		480 Leu
	Cys	Ala	Asp		485 Tyr	Phe	Gly	Asp		490 Phe	Gly	Glu	Arg	Gly	495 Pro	Val
	Arg	Pro		500 Gln	Pro	Cys	Gln		505 Asn	Asn	Asn	Val		510 Pro	Ser	Ala
	Ser	_	515 Asn	Cys	Asp	Arg		520 Thr	Gly	Arg	Cys		525 Lys	Cys	Ile	His
		530 Thr	Ala	Gly	Val		535 Cys	Asp	Gln	Cys	Lys 555	540 Ala	Gly	Tyr	Tyr	Gly 560
	545 Asp	Pro	Leu	Ala	Pro 565	550 Asn	Pro	Ala	Asp	Lys 570		Arg	Ala	Суѕ	Asn 575	
	Asn	Pro	Val	Gly 580		Glu	Pro	Val	Glu 585		Arg	Ser	Asp	Gly 590		Cys
	Val	Cys	Lys 595		Gly	Phe	Gly	Gly 600		Ser	Cys	Glu	His 605	Ala	Ala	Leu
	Thr	Ser 610		Pro	Ala	Cys	Tyr 615		Gln	Val	Lys	Val 620		Met	Asp	Gln
	Phe 625		Gln	Gln	Leu	Gln 630		Leu	Glu	Ala	Leu 635		Ser	Lys	Ala	Gln 640
		Gly	Ala	Val	Pro 645		Ala	Glu	Leu	Glu 650		Arg	Met	Gln	Gln 655	
	Glu	Gln	Ala	Leu 660		Asp	Ile	Leu	Arg 665		Ala	Gln	Ile	Ser 670	Gln	Asp
	Ala	Val	Arg 675		Phe	Asn	Leu	Arg 680		Ala	Lys	Ala	Arg 685	Thr	Gln	Glu
	Asn	Ser 690	Tyr	Arg	Asp	Arg	Leu 695	Asp	Asp	Leu	Lys	Met 700	Thr	Val	Glu	Arg
	Val 705		Ala	Leu	Gly	Ser 710	Gln	Tyr	Gln	Asn	Gln 715	Val	Gln	Asp	Thr	Arg 720
	Arg	Leu	Ile	Thr	Gln 725	Met	Arg	Leu	Ser	Leu 730	Glu	Glu	Ser	Glu	Ala 735	Ser
	Leu	Gln	Asn	Thr 740	Asn	Ile	Pro	Pro	Ser 745	Glu	His	Tyr	Val	Gly 750	Pro	Asn
	_		755					760					765	Asp		
	Val	Gln 770	Ser	Ala	Ser	Asn	Met 775	Glu	Gln	Leu	Ala	Lys 780	Glu	Thr	Gln	Glu
	785					790					795			Gln		800
					805					810				Leu	815	
	_			820		_			825					Arg 830		
			835					840					845	Ser		
	Leu	Leu 850	Asn	Ser	Val	Ser	Gln 855	Ile	Gln	Gly	Val	Asn 860	Asp	Gln	Ser	Leu

```
Gln Val Glu Ala Lys Arg Leu Arg Gln Lys Ala Asp Ser Leu Ser Asn
                  870
Arg Val Thr Lys His Met Asp Glu Phe Lys His Val Gln Ser Asn Leu
                                 890
Gly Asn Trp Glu Glu Glu Thr Arg Gln Leu Leu Gln Asn Gly Lys Asn
                             905
Gly Arg Gln Thr Ser Asp Gln Leu Leu Ser Arg Ala Asn Leu Ala Lys
               920
Ser Arg Ala Gln Glu Ala Leu Ser Met Gly Asn Ala Thr Phe Tyr Glu
           935
Val Glu Asn Ile Leu Lys Asn Leu Arg Glu Phe Asp Leu Gln Val Gly
                                    955
                  950
Asp Lys Arg Ala Glu Ala Glu Glu Ala Met Lys Arg Leu Ser Tyr Ile
                                970
              965
Ser Gln Lys Val Ala Gly Ala Ser Asp Lys Thr Lys Gln Ala Glu Ala
                             985
           980
Ala Leu Gly Ser Ala Ala Ala Asp Ala Gln Arg Ala Lys Asn Ala Ala
                         1000
                                            1005
Arg Glu Ala Leu Glu Ile Ser Gly Lys Ile Glu Gln Glu Ile Gly Gly
                     1015
                                        1020
Leu Asn Leu Glu Ala Asn Val Thr Ala Asp Gly Ala Leu Ala Met Glu
                 1030
                          1035
Lys Gly Leu Ala Thr Leu Lys Ser Glu Met Arg Glu Val Glu Gly Glu
                      1050
             1045
Leu Ser Arg Lys Glu Gln Glu Phe Asp Met Asp Met Asp Ala Val Gln
                            1065
          1060
Met Val Ile Ala Glu Ala Gln Arg Val Glu Asn Arg Ala Lys Asn Ala
                                1085
                        1080
Gly Val Thr Ile Gln Asp Thr Leu Asn Thr Leu Asp Gly Ile Leu His
   1090 . 1095
                                        1100
Leu Ile Asp Gln Pro Gly Ser Val Asp Glu Glu Arg Leu Ile Leu Leu
                                    1115
                  1110
Glu Gln Lys Leu Phe Arg Ala Lys Thr Gln Ile Asn Ser Gln Leu Arg
                                1130
              1125
Pro Leu Met Ser Glu Leu Glu Glu Arg Ala His Arg Gln Lys Gly His
                            1145
          1140
Leu Arg Phe Leu Glu Thr Ser Ile Asp Gly Ile Leu Ala Asp Val Lys
                        1160
                                           1165
Asn Leu Glu Asn Ile Arg Asp Asn Leu Pro Pro Gly Cys Tyr Asn Thr
                    1175
Gln Ala Leu Glu Gln Gln
1185
<210> 3
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 3
atgcctgcgc tctggctcag
```

<210> 4 <211> 20 <212> DNA

<213> Artificial Sequence	
<220>	
<223> primer	
<400> 4	20
tgtggcagct ggcggaatgc	20
<210> 5	
<211> 20	
<212> DNA <213> Artificial Sequence	
22132 Artificial Sequence	
<220>	
<223> primer	
<400> 5	
gactccaagt gtgactgtga	20
<210> 6	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> primer	
<400> 6	
tagctcacct gttgattccc	20
<210> 7	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> primer	
<400> 7	
cctgtctatt ttgtagctcc	20
<210> 8	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> primer	
<400> 8	
cagetgaace cattgegaca	20
<210> 9	
<211> 20	
<212> DNA	
<213> Artificial Sequence	

·<220>	
<223> primer	
<400> 9	
gagtatcgga ggttactgcg	20
<210> 10	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> primer	
<400> 10	
gacactccac aggctccgag	20
<210> 11	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> primer	
•	
<400> 11	
cgctgtgagc tctgtgctga	20
<210> 12	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> primer	
<400> 12	
ctccaggatc tggagctgct	20
<210> 13	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> primer	
· · · · · · · · · · · · · · · · · · ·	
<400> 13	
gacaagtgtc gagcttgcaa	20
<210> 14	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
•	
<220>	
<223> primer	
-	

<400> 14 tcatcatgag gtcatccagg	20
<210> 15 <211> 20	
<212> DNA	
<213> Artificial Sequence	•
<220> <223> primer	
<400> 15	20
gagagaagcc cagatttcac	20
<210> 16	
<211> 20	
<212> DNA <213> Artificial Sequence	
<2213> Altilitat Sequence	
<220>	
<223> primer	
<400> 16	
gettecatgt eggtttgegt	20
<210> 17	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> primer	
<400> 17	
cagccagtaa catggagcaa	20
<210> 18	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> primer	
<400> 18	
gtctgtctcc cattctttcc	20
<210> 19	
<211> 20	
<212> DNA <213> Artificial Sequence	
<220>	
<223> primer	
<400> 19	

į.	n .		
•			
	•		
•	ggatgagttc aagcacgtgc		20
	<210> 20		
	<211> 20		
	<212> DNA		
	<213> Artificial Sequence		
	<220>		
	<223> primer		
	(225) primer		
	<400> 20		
			20
	acagetetee ttecaettet		20
	<210> 21		
	<211> 20		
	<212> DNA		
	<213> Artificial Sequence		
	<220>		
	<223> primer		
	-		
	<400> 21		
	caagacgaag caagcagaag		20
	<210> 22		
	<211> 20		
	<212> DNA		
	<213> Artificial Sequence		
	2213) Altificial Sequence		
	222		
	<220>		
	<223> primer		
	<400> 22		
	ggctgttgat ctgagtcttg		20
	<210> 23	•	
	<211> 20		
	<212> DNA		
	<213> Artificial Sequence		
	<220>		
	<223> primer		
	<400> 23		
	gtgagtcaca ccctgaaaca		20
	gegageeaca eeeegaaaca		
	<210> 24		
•	<210> 24 <211> 22		
	<212> DNA		
	<213> Artificial Sequence		
	<220>		
	<223> primer		
	<400> 24		
	gagttacaat tqcagggtaa ac		22

```
`<210> 25
<211> 36
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 25
                                                                           36
ggccatgcgt agactcttaa ttttttttt ttttt
<210> 26
<211> 19
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 26
                                                                           19
gcagaggccc aaagagttg
<210> 27
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 27
                                                                           20
ggccatgcgt agactcttaa
<210> 28
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 28
                                                                           20
cctggcagtg tggatgaaga
<210> 29
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 29
                                                                           20
tgttactcag gggatgagaa
<210> 30
<211> 19
```

```
<213> Artificial Sequence
<220>
<223> primer
<400> 30
ctgggggcag ttattgcac
<210> 31
<211> 1193
<212> PRT
<213> human
<220>
<223>
<400> 31
Met Pro Ala Leu Trp Leu Gly Cys Cys Leu Cys Phe Ser Leu Leu Leu
Pro Ala Ala Arg Ala Thr Ser Arg Arg Glu Val Cys Asp Cys Asn Gly
Lys Ser Arg Gln Cys Ile Phe Asp Arg Glu Leu His Arg Gln Thr Gly
                            40
Asn Gly Phe Arg Cys Leu Asn Cys Asn Asp Asn Thr Asp Gly Ile His
                        55
Cys Glu Lys Cys Lys Asn Gly Phe Tyr Arg His Arg Glu Arg Asp Arg
                   70
                                        75
Cys Leu Pro Cys Asn Cys Asn Ser Lys Gly Ser Leu Ser Ala Arg Cys
Asp Asn Ser Gly Arg Cys Ser Cys Lys Pro Gly Val Thr Gly Ala Arg
                                105
Cys Asp Arg Cys Leu Pro Gly Phe His Met Leu Thr Asp Ala Gly Cys
                            120
Thr Gln Asp Gln Arg Leu Leu Asp Ser Lys Cys Asp Cys Asp Pro Ala
                       135
Gly Ile Ala Gly Pro Cys Asp Ala Gly Arg Cys Val Cys Lys Pro Ala
                    150
                                        155
Val Thr Gly Glu Arg Cys Asp Arg Cys Arg Ser Gly Tyr Tyr Asn Leu
                                    170
               165
Asp Gly Gly Asn Pro Glu Gly Cys Thr Gln Cys Phe Cys Tyr Gly His
                                                    190
                                185
Ser Ala Ser Cys Arg Ser Ser Ala Glu Tyr Ser Val His Lys Ile Thr
                            200
Ser Thr Phe His Gln Asp Val Asp Gly Trp Lys Ala Val Gln Arg Asn
                        215
Gly Ser Pro Ala Lys Leu Gln Trp Ser Gln Arg His Gln Asp Val Phe
                    230
                                        235
Ser Ser Ala Gln Arg Leu Asp Pro Val Tyr Phe Val Ala Pro Ala Lys
                                   250
Phe Leu Gly Asn Gln Gln Val Ser Tyr Gly Gln Ser Leu Ser Phe Asp
            260
                                265
Tyr Arg Val Asp Arg Gly Gly Arg His Pro Ser Ala His Asp Val Ile
                            280
       275
Leu Glu Gly Ala Gly Leu Arg Ile Thr Ala Pro Leu Met Pro Leu Gly
                                            300
                        295
Lys Thr Leu Pro Cys Gly Leu Thr Lys Thr Tyr Thr Phe Arg Leu Asn
```

19

<212> DNA

ı																
, .							٠								•	
•	3				•											
	305					310					315					320
		His	Pro	Ser	Asn 325		Trp	Ser	Pro	Gln 330		Ser	Tyr	Phe	Glu 335	Tyr
	Arg	Arg	Leu	Leu 340	Arg	Asn	Leu	Thr	Ala 345	Leu	Arg	Ile	Arg	Ala 350	Thr	Tyr
,	Gly	Glu	Tyr 355	Ser	Thr	Gly	Tyr	Ile 360	Asp	Asn	Val	Thr	Leu 365	Ile	Ser	Ala
	_	370	Val		-		375					380				
	385		Gly	_	_	390					395					400
	_	_	Asp		405					410					415	
		_	Gln	420					425					430		
		-	Asp 435					440					445			
		450	Asn				455					460				
	465		Gly			470					475					480
	-		Asn	_	485					490					495	
			Gly	500					505					510		
		_	Gln 515 Cys					520					525			
	-	530	Gly	_			535					540				
	545		Ala		_	550					555					560
					565					570					575	Val
			_	580					585					590		Ser
	_	•	595 Ala					600					605			
	_	610		_	_		615					620				Gly
	625 Asp	Gly	Val	Val	Pro	630 Asp	Thr	Glu	Leu	Glu	635 Gly	Arg	Met	Gln	Gln	640 Ala
1	Glu	Gln	Ala	Leu	645 Gln	Asp	Ile	Leu	Arg	650 Asp	Ala	Gln	Ile	Ser	655 Glu	Gly
	Ala	Ser	Arg	660 Ser	Leu	Gly	Leu	Gln	665 Leu	Ala	Lys	Val	Arg	670 Ser	Gln	Glu
	Asn		675 Tyr	Gln	Ser	Arg		680 Asp	Asp	Leu	Lys		685 Thr	Val	Glu	Arg
		690 Arg	Ala	Leu	Gly		695 Gln	Tyr	Gln	Asn		700 Val	Arg	Asp	Thr	His
	705 Arg	Leu	Ile	Thr		710 Met	Gln	Leu	Ser		715 Ala	Glu	Ser	Glu		720 Ser
	Leu	Gly	Asn		725 Asn	Ile	Pro	Ala	Ser	730 Asp	His	Tyr	Val	Gly 750	735 Pro	Asn
	Gly	Phe	Lys 755	740 Ser	Leu	Ala	Gln	Glu 760		Thr	Arg	Leu	Ala 765		Ser	His

Val Glu Ser Ala Ser Asn Met Glu Gln Leu Thr Arg Glu Thr Glu Asp Tyr Ser Lys Gln Ala Leu Ser Leu Val Arg Lys Ala Leu His Glu Gly Val Gly Ser Gly Ser Gly Ser Pro Asp Gly Ala Val Val Gln Gly Leu Val Glu Lys Leu Glu Lys Thr Lys Ser Leu Ala Gln Gln Leu Thr Arg . 830 Glu Ala Thr Gln Ala Glu Ile Glu Ala Asp Arg Ser Tyr Gln His Ser Leu Arg Leu Leu Asp Ser Val Ser Arg Leu Gln Gly Val Ser Asp Gln Ser Phe Gln Val Glu Glu Ala Lys Arg Ile Lys Gln Lys Ala Asp Ser Leu Ser Thr Leu Val Thr Arg His Met Asp Glu Phe Lys Arg Thr Gln Lys Asn Leu Gly Asn Trp Lys Glu Glu Ala Gln Gln Leu Leu Gln Asn Gly Lys Ser Gly Arg Glu Lys Ser Asp Gln Leu Leu Ser Arg Ala Asn Leu Ala Lys Ser Arg Ala Gln Glu Ala Leu Ser Met Gly Asn Ala Thr Phe Tyr Glu Val Glu Ser Ile Leu Lys Asn Leu Arg Glu Phe Asp Leu Gln Val Asp Asn Arg Lys Ala Glu Ala Glu Glu Ala Met Lys Arg Leu Ser Tyr Ile Ser Gln Lys Val Ser Asp Ala Ser Asp Lys Thr Gln Gln Ala Glu Arg Ala Leu Gly Ser Ala Ala Ala Asp Ala Gln Arg Ala Lys Asn Gly Ala Gly Glu Ala Leu Glu Ile Ser Ser Glu Ile Glu Gln Glu Ile Gly Ser Leu Asn Leu Glu Ala Asn Val Thr Ala Asp Gly Ala Leu Ala Met Glu Lys Gly Leu Ala Ser Leu Lys Ser Glu Met Arg Glu Val Glu Gly Glu Leu Glu Arg Lys Glu Leu Glu Phe Asp Thr Asn Met Asp Ala Val Gln Met Val Ile Thr Glu Ala Gln Lys Val Asp Thr Arg Ala Lys Asn Ala Gly Val Thr Ile Gln Asp Thr Leu Asn Thr Leu Asp Gly Leu Leu His Leu Met Asp Gln Pro Leu Ser Val Asp Glu Glu Gly Leu Val Leu Leu Glu Gln Lys Leu Ser Arg Ala Lys Thr Gln Ile Asn Ser Gln Leu Arg Pro Met Met Ser Glu Leu Glu Glu Arg Ala Arg Gln Gln Arg Gly His Leu His Leu Leu Glu Thr Ser Ile Asp Gly Ile Leu Ala Asp Val Lys Asn Leu Glu Asn Ile Arg Asp Asn Leu Pro Pro Gly Cys Tyr Asn Thr Gln Ala Leu Glu Gln Gln 

<212> PRT
<213> murine
<220>
<223>
<400> 32
Met Pro Ala
1
Pro Ala Ser
Lys Ser Arg

<400> 32 Met Pro Ala Leu Trp Leu Ser Cys Cys Leu Gly Val Ala Leu Leu Leu 10 Pro Ala Ser Gln Ala Thr Ser Arg Glu Val Cys Asp Cys Asn Gly 25 Lys Ser Arg Gln Cys Val Phe Asp Gln Glu Leu His Arg Gln Ala Gly 40 Ser Gly Phe Arg Cys Leu Asn Cys Asn Asp Asn Thr Ala Gly Val His Cys Glu Arg Ser Arg Glu Gly Phe Tyr Gln His Gln Ser Lys Ser Arg Cys Leu Pro Cys Asn Cys His Ser Lys Gly Ser Leu Ser Ala Gly Cys 90 Asp Asn Ser Gly Gln Cys Arg Cys Lys Pro Gly Val Thr Gly Gln Arg 105 Cys Asp Gln Cys Gln Pro Gly Phe His Met Leu Thr Asp Ala Gly Cys 125 120 Thr Arg Asp Gln Gly Gln Leu Asp Ser Lys Cys Asp Cys Asp Pro Ala 135 140 Gly Ile Ser Gly Pro Cys Asp Ser Gly Arg Cys Val Cys Lys Pro Ala 150 155 Val Thr Gly Glu Arg Cys Asp Arg Cys Arg Pro Arg Asp Tyr His Leu 170 165 Asp Arg Ala Asn Pro Glu Gly Cys Thr Gln Cys Phe Cys Tyr Gly His 185 Ser Ala Ser Cys His Ala Ser Ala Asp Phe Ser Val His Lys Ile Thr 200 205 Ser Thr Phe Ser Gln Asp Val Asp Gly Trp Lys Ala Val Gln Arg Asn 215 Gly Ala Pro Ala Lys Leu His Trp Ser Gln Arg His Arg Asp Val Phe 230 235 Ser Ser Ala Arg Arg Ser Asp Pro Val Tyr Phe Val Ala Pro Ala Lys 250 245 Phe Leu Gly Asn Gln Gln Val Ser Tyr Gly Gln Ser Leu Ser Phe Asp 265 Tyr Arg Val Asp Arg Gly Gly Arg Gln Pro Ser Ala Tyr Asp Val Ile 280 Leu Glu Gly Ala Gly Leu Gln Ile Arg Ala Pro Leu Met Ala Pro Gly 295 Lys Thr Leu Pro Cys Gly Ile Thr Lys Thr Tyr Thr Phe Arg Leu Asn 315 310 Glu His Pro Ser Ser His Trp Ser Pro Gln Leu Ser Tyr Phe Glu Tyr 330 325 Arg Arg Leu Leu Arg Asn Leu Thr Ala Leu Leu Met Ile Arg Ala Thr 345 350 Tyr Gly Glu Tyr Ser Thr Gly Tyr Ile Asp Asn Val Thr Leu Val Ser 360 365 Ala Arg Pro Val Leu Gly Ala Pro Ala Pro Trp Val Glu Arg Cys Val 375 380 Cys Leu Leu Gly Tyr Lys Gly Gln Phe Cys Gln Glu Cys Ala Ser Gly 395 390

Tyr Lys Arg Asp Ser Ala Arg Leu Gly Ala Phe Gly Ala Cys Val Pro 410 Cys Asn Cys Gln Gly Glu Gly Ala Cys Asp Pro Asp Thr Gly Asp Cys 425 Tyr Ser Gly Asp Glu Asn Pro Asp Ile Glu Cys Ala Asp Cys Pro Ile 440 Gly Phe Tyr Asn Asp Pro His Asp Pro Arg Ser Cys Lys Pro Cys Pro 455 Cys His Asn Gly Phe Ser Cys Ser Val Met Pro Glu Thr Glu Glu Val 470 475 Val Cys Asn Asn Cys Pro Pro Gly Val Thr Gly Ala Arg Cys Glu Leu 485 490 Cys Ala Asp Gly Phe Phe Gly Asp Pro Phe Gly Glu His Gly Pro Val 505 510 Arg Pro Cys Gln Arg Cys Gln Cys Asn Asn Asn Val Asp Pro Asn Ala 520 Ser Gly Asn Cys Asp Gln Leu Thr Gly Arg Cys Leu Lys Cys Ile Tyr 535 Asn Thr Ala Gly Val Tyr Cys Asp Gln Cys Lys Ala Gly Tyr Phe Gly 555 550 Asp Pro Leu Ala Pro Asn Pro Ala Asp Lys Cys Arg Ala Cys Asn Cys 570 565 Ser Pro Met Gly Ala Glu Pro Gly Glu Cys Arg Gly Asp Gly Ser Cys 585 Val Cys Lys Pro Gly Phe Gly Ala Phe Asn Cys Asp His Ala Ala Leu 605 600 Thr Ser Cys Pro Ala Cys Tyr Asn Gln Val Lys Ile Gln Met Asp Gln 615 Phe Thr Gln Gln Leu Gln Ser Leu Glu Ala Leu Val Ser Lys Ala Gln 635 630 Gly Gly Gly Gly Gly Thr Val Pro Val Gln Leu Glu Gly Arg Ile 650 645 Glu Gln Ala Glu Gln Ala Leu Gln Asp Ile Leu Gly Glu Ala Gln Ile 665 Ser Glu Gly Ala Met Arg Ala Val Ala Val Arg Leu Ala Lys Ala Arg 680 Ser Gln Glu Asn Asp Tyr Lys Thr Arg Leu Asp Asp Leu Lys Met Thr 700 695 Ala Glu Arg Ile Arg Ala Leu Gly Ser Gln His Gln Asn Arg Val Gln 715 710 Asp Thr Ser Arg Leu Ile Ser Gln Met Arg Leu Ser Leu Ala Gly Ser 730 725 Glu Ala Leu Leu Glu Asn Thr Asn Ile His Ser Ser Glu His Tyr Val 745 Gly Pro Asn Asp Phe Lys Ser Leu Ala Gln Glu Ala Thr Arg Lys Ala 760 Asp Ser His Ala Glu Ser Ala Asn Ala Met Lys Gln Leu Ala Arg Glu 775 780 Thr Glu Asp Tyr Ser Lys Gln Ala Leu Ser Leu Ala Arg Lys Leu Leu 795 790 Ser Gly Gly Gly Ser Gly Ser Trp Asp Ser Ser Val Val Gln Gly 810 Leu Met Gly Lys Leu Glu Lys Thr Lys Ser Leu Ser Gln Gln Leu Ser 820 825 Leu Glu Gly Thr Gln Ala Asp Ile Glu Ala Asp Arg Ser Tyr Gln His 840 Ser Leu Arq Leu Leu Asp Ser Ala Ser Gln Leu Gln Gly Val Ser Asp

Leu Ser Phe Gln Val Glu Ala Lys Arg Ile Arg Gln Lys Ala Asp Ser Leu Ser Asn Leu Val Thr Arg Gln Thr Asp Ala Phe Thr Arg Val Arg Asn Asn Leu Gly Asn Trp Glu Lys Glu Thr Arg Gln Leu Leu Gln Thr Gly Lys Asp Arg Arg Gln Thr Ser Asp Gln Leu Leu Ser Arg Ala Asn Leu Ala Lys Asn Arg Ala Gln Glu Ala Leu Ser Met Gly Asn Ala Thr Phe Tyr Glu Val Glu Asn Ile Leu Lys Asn Leu Arg Glu Phe Asp Leu Gln Val Glu Asp Arg Lys Ala Glu Ala Glu Glu Ala Met Lys Arg Leu Ser Ser Ile Ser Gln Lys Val Ala Asp Ala Ser Asp Lys Thr Gln Gln Ala Glu Thr Ala Leu Gly Ser Ala Thr Ala Asp Thr Gln Arg Ala Lys Asn Ala Ala Arg Glu Ala Leu Glu Ile Ser Ser Glu Ile Glu Leu Glu Ile Gly Ser Leu Asn Leu Glu Ala Asn Val Thr Ala Asp Gly Ala Leu Ala Met Glu Lys Gly Thr Ala Thr Leu Lys Ser Glu Met Arg Glu Met Ile Glu Leu Ala Arg Lys Glu Leu Glu Phe Asp Thr Asp Lys Asp Thr Val Gln Leu Val Ile Thr Glu Ala Gln Gln Ala Asp Ala Arg Ala Thr Ser Ala Gly Val Thr Ile Gln Asp Thr Leu Asn Thr Leu Asp Gly Ile Leu His Leu Ile Asp Gln Pro Gly Ser Val Asp Glu Glu Gly Met Met Leu Leu Glu Gln Gly Leu Phe Gln Ala Lys Thr Gln Ile Asn Ser Arg Leu Arg Pro Leu Met Ser Asp Leu Glu Glu Arg Val Arg Arg Gln Arg Asn His Leu His Leu Leu Glu Thr Ser Ile Asp Gly Ile Leu Ala Asp Val Lys Asn Leu Glu Asn Ile Arg Asp Asn Leu Pro Pro Gly Cys Tyr Asn Thr Gln Ala Leu Glu Gln Gln